

The Virtual Institute of Microbial Stress and Survival Rapid Deduction of Stress Response Pathways in Metal/Radionuclide Reducing Ba

Applied Environmental Microbiology Core





Rapid Deduction of Stress Response Pathways in Metal/Radionuclide Reducing Bacteria

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Overview







AEMC Mission

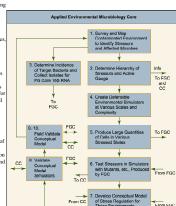
AEMC is the source of environmental data and samples that determine the stressors that will be studied, provides the environments for growing the organisms to be tested, simulates stressed environments, and verifies the conceptual models to determine how these stress regulatory pathways control the biogeochemistry of contaminated sites

Main Goals

- Develop criteria for monitoring the integrity (health) and altering the trajectory of an environmental biological system (process control).
- More complete understanding of the diversity and environmental context of stress response.

Specific Aims

- Survey and map DOE sites contaminated by metals and radionuclides using chemical and molecular/microbiological parameters to determine major microbial populations and potential stressors for Desulfovibrio vulgaris, Geobacter metallireducens, and Shewanella oneidensis MR1.
- Determine the rank priority of these stressors in terms of their ability to
 affect metal/radionuclide bioreduction by either direct or indirect processes
 and to establish the normal active range of the stressor in
 metal/radionuclide contaminated environments.
- Determine the incidence and activity of the three target bacteria, and closely related relatives, in the test metal/radionuclide contaminated environments and collect isolates for analysis by the Functional Genomic Core for comparison using 165 RNA profiling.
- Create defensible environmental simulators that can replicate key features
 of field site chemical and biological structure to mimic stress conditions for
 single populations and later for microbial communities (chemostats to soil
 columns from 10 μm to 1 m size systems).
- Provide large quantities of cells in various stress states for the Functional Genomics Core's physiological monitoring facility, and molecular interaction studies.
- Provide environmental simulators for testing stressor effects on mutants, large insert clones, expression analysis, etc., for elucidating critical parts of the stress regulation pathway.
- Develop testable conceptual models of stress regulatory pathways based on results of the Computational Core that could predict natural attenuation and suggest biostimulatory strategies for immobilization of metals and radionuclides at DOE contaminated sites.
- Test conceptual models of stress regulatory pathways and effects on contaminate site biogeochemistry using competent soil columns with different levels of complexity over the active range of the stressors
- Validate conceptual models using field tests at contaminated sites that utilize specific functional gene arrays developed from the stress regulator pathways.
- Alter field conditions or test along gradients to verify stress regulatory
 model efficiency for predicting natural attenuation or suggesting
 biostimulatory strategies for immobilization of metals and radionuclides



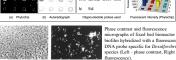
Environmental Simulator Stress Analyses

Growth and recovery of stressed single populations under simple conditions
Identification and recovery of stressresponsive populations from complex communities

Radiomicroarray analysis
 Recover of stress responsive
 populations via flow cytometry
 Flow cytometric sorting of FISH-labeled populations

Expresssion analysis of RNA extraction from sediment columns and reactor systems
 Synchrotron FTIR direct analysis of stress changes in living cells
 Direct stress and community comparison with PLFA and metabolite.

analyses



Phylogenetic affiliation of bioreactor isolate PT2. This organism comprised a significant fraction of the Pradigoritor, openulation identifies in the population described in the production of the production

UniOT Sper152(5) Actitis Nacrillis

Reactor System Development

Rational

- Analogous to saturated soil/sediment systems in which the microbiota coloniz
- surfaces as biofilms

 Growth as biofilms is the more common state for microorganisms in the
- Fluidized bed reactor systems provide for experimental control, replication biomass, and ease of sampling
- relevant state for most environmental populations

 Biofilm-associated populations are not washed out (of the reactor system) following imposition of stress, providing for greater range of stress-respondence.
- Past reactor studies have shown that Desulfovibrio spp. and Geobacter-lik populations will colonize and be retained within these systems.
- populations will colonize and be retained within these systems

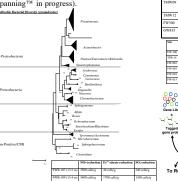
 Chemostat Studies
- Cultural isolation of organisms for comparative "Benchmarks'
 Cultural isolation of antisymmetral complex for progress incompeting
- Collection of environmental samples for reactor inoculation
 Comparability of reactor and field site community structure



Synthetic soil aggregate column design [6] (1) Homogeneous soil pack, (2) plastic column, (3) piston for soil extrusion, (6) boundary reservoir for Crt/VI) solution (5) Pt redox electrodes, and (6) plug.

Chemical & Molecular Microbiological Mapping of Field Sites

- Survey of existing data at the NABIR Field Research Center at Oak Ridge - Draft report completed.
- Additional sediment and soil sampling for: physical/chemical stressors, wild isolates of target bacteria, and TRFLP and clonal population mapping (Biopanning TM in progress).



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	TPS-16	47%	244	7.1	73	
	FW-003	321	34	3.0	3.9	
	PW-005	216	47	3.8	6.3	
	I-W-010	115	27	3.1	43	
	PW-015	229	61	4.3	9.1	
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Validation of Conceptual Models with Simulators and Field Tests

This research element will not begin until the 3rd year after pathways have been identified and tested.

Experimental Core Facilities

Environmental Molecular Microbiology Facility (ORNL, LBNL, U. Wash., Diversa Inc.): Affymetrix microarray system, PLFA analysis lab, Radio-microarray lab, low capacity DNA microarrayer, flow cytometer, Laser confocal microscope lab, Synchrotron FTIR beamline (Advanced Light Source), clone libraries from NABIR FRC, Microarrays for Shewanella

Environmental Simulation and Culture Facility (LBNL, U. Wash., Diversa Inc., U. Missouri): Anaerobic chambers with built-in microscopes and incubators, instrumented soil columns (various sizes). Ultracold freezers & liquid nitrogen archiving, Biopanning ^{pat}, Chemostats (Fermentis), Batch Fermentors, Fixed bed reactors, Fluidized bed reactors, radiological control areas for microbiological sample prep and incubation

